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DEFINITION Tetraodon nigroviridis genome survey sequence 17 end of clone
LOCUS A2311.1A library B from Tetraodon nigroviridis, genomic survey
sequence.
VERSION A2311.1A
KEYWORDS GSS, genome survey sequence,
SOURCE Tetraodon nigroviridis,
ORGANISM Tetraodon nigroviridis
Eukaryota; Metazoa; Chordata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Euteleostei; Reueteleostei;
Actinopterygii; Acanthopterygii; Perciformes; Tetraodontiformes;
Tetraodontidae; Tetraodonti.
REFERENCE 1. (bases 1 to 1101)
Kobayashi T, Miller H, Miller H, Dastler C, Fuzumori O, Fisher C,
Homenick J, Billault A, Quetier F, Saurin W, Bernot A, and
Weissenbach J.
TITLE Characterization and repeat analysis of the compact genome of the
freshwater pufferfish Tetraodon nigroviridis
JOURNAL Republished
AUTHORS 2. (bases 1 to 1101)
Kobayashi T, Miller H, Miller H, Dastler C, Fuzumori O, Fisher C,
Homenick J, Billault A, Quetier F, Saurin W, Bernot A, and
Weissenbach J.
AUTHORS 3. (bases 1 to 1101)
Kobayashi T, Miller H, Miller H, Dastler C, Fuzumori O, Fisher C,
Homenick J, Billault A, Quetier F, Saurin W, Bernot A, and
Weissenbach J.
TITLE Human genome number estimate provided by genome wide analysis using
Tetraodon nigroviridis DNA sequence
JOURNAL Republished
AUTHORS 4. (bases 1 to 1101)
Kobayashi T, Miller H, Miller H, Dastler C, Fuzumori O, Fisher C,
Homenick J, Billault A, Quetier F, Saurin W, Bernot A, and
Weissenbach J.
TITLE Direct Submission
JOURNAL Published in Apr 2002 to the PDB, GenBank, and EMBL
COMMENT This sequence is a single read and was generated as part of a large
scale clone-end sequencing project of the Tetraodon nigroviridis
genome. For more information, please take a look at
http://www.genoscope.cns.fr/708a-15.
FEATURES
Source Location/Qualifiers
1..1101
Organism: "Tetraodon nigroviridis"
/plasmid: "taxon:99883"
/db_xref: "taxon:99883"
/clone_lib: "B"
/note: "Genoscope sequence ID : COAB02PAB0602 end : T7"
BASE COUNT 326 a 154 c 172 g 378 t 72 others
DEFINITION 15
LOCUS CNS016E2 1204 bp DNA linear GSS 26-JUL-1999
DEFINITION Drosophila melanogaster genome survey sequence T7 end of BAC
BACN15A12 of DrosBAC library from Drosophila melanogaster (fruit
fly), genomic survey sequence.
ACCESSION A0106628.1 GI:5622852
VERSION A0106628.1
KEYWORDS GSS
SOURCE Drosophila melanogaster
ORGANISM Drosophila melanogaster
Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
Plecoptera; Megoptera; Megoptera; Diptera; Arthropoda;
Muscomorpha; Ephyraeidae; Drosophilidae; Drosophila.
REFERENCE 1. (bases 1 to 1204)
Genoscope.
AUTHORS Direct Submission
JOURNAL Submitted (23-JUL-1999) Genoscope - Centre National de Sequencage :
BP 191 91006 EVRY cedex - FRANCE (E-mail : seq@genoscope.cns.fr
- Web : www.genoscope.cns.fr)
COMMENT Determination of this BAC-end sequence was carried out as part of a
collaboration with the European Drosophila Genome Project (EDGP) -
http://www.edgp.sbi.ac.uk. This Drosophila melanogaster BAC
library (Dros BAC) was made by Alain Billaut at CEPH (Centre
d'Etude du Polymorphisme Humain) with funding provided by a MC
project grant. The DNA was prepared from embryos by Alain Bucheton
and Genevieve Lefebvre. It has been constructed in the vector
pBeloBAC11.
FEATURES
Source Location/Qualifiers
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/plasmid: "pBeloBAC11"
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/clone_lib: "DrosBAC"
/clone: "BACN15A12"
/note: "end : T7"
BASE COUNT 298 a 172 c 106 g 312 t 312 others
DEFINITION 1248
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DEFINITION Drosophila melanogaster genome survey sequence T7 end of BAC
BACN15A12 of DrosBAC library from Drosophila melanogaster (fruit
fly), genomic survey sequence.
ACCESSION A0106628.1 GI:5622852
VERSION A0106628.1
KEYWORDS GSS
SOURCE Drosophila melanogaster
ORGANISM Drosophila melanogaster
Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
Plecoptera; Megoptera; Megoptera; Diptera; Arthropoda;
Muscomorpha; Ephyraeidae; Drosophilidae; Drosophila.
REFERENCE 1. (bases 1 to 1204)
Genoscope.
AUTHORS Direct Submission
JOURNAL Submitted (23-JUL-1999) Genoscope - Centre National de Sequencage :
BP 191 91006 EVRY cedex - FRANCE (E-mail : seq@genoscope.cns.fr
- Web : www.genoscope.cns.fr)
COMMENT Determination of this BAC-end sequence was carried out as part of a
collaboration with the European Drosophila Genome Project (EDGP) -
http://www.edgp.sbi.ac.uk. This Drosophila melanogaster BAC
library (Dros BAC) was made by Alain Billaut at CEPH (Centre
d'Etude du Polymorphisme Humain) with funding provided by a MC
project grant. The DNA was prepared from embryos by Alain Bucheton
and Genevieve Lefebvre. It has been constructed in the vector
pBeloBAC11.
FEATURES
Source Location/Qualifiers
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Organism: "Drosophila melanogaster"
/plasmid: "pBeloBAC11"
/db_xref: "taxon:7227"
/clone_lib: "DrosBAC"
/clone: "BACN15A12"
/note: "end : T7"
BASE COUNT 298 a 172 c 106 g 312 t 312 others

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Wed Jun 5 09:42:34 2002

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